

GACGTAGGATCCAAGCGTTATCCGGAATTACTGGGCGTAAAGGGCGTGTA GGAGGTTGGGCAAGTCGGCCATGAAAGCTCCCGGCTCAAC

TAXONOMY: Assigned Greengenes taxonomy & most common taxonomy of 100% ID matches to RDP.

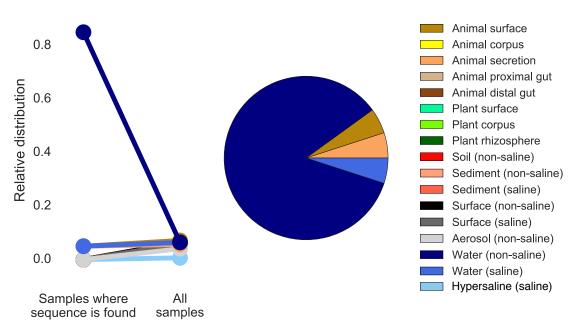
Greengenes lineage:	k_Bacteria; p_Chloroflexi; c_Anaerolineae; o_H39; f_; g_; s_
RDP lineage:	d_Bacteria; k_; p_Candidatus Saccharibacteria; c_; o_; f_; g_Saccharib- acteria_genera_incertae_sedis (1/1)
RDP species:	metal-contaminated soil (1/1)

PREVALENCE: Found in 1.00% of samples, rank #7840 out of 153302 tag sequences.

ABUNDANCE: Composes 0.117% of observations, rank #92 out of 153302 tag sequences.

METHODS: Amplicon PCR with 16S rRNA V4 primers 515f–806rB. Sequencing with Illumina HiSeq 90/100-bp or MiSeq 150-bp single reads. Sequences error-checked with Deblur, trimmed to 90 bp & rarefied to 5000 obs./sample. Showing 2000 samples evenly distributed by habitat & study.

PREVALENCE BY HABITAT:



ABUNDANCE BY ENVIRONMENTAL PARAMETERS:

